EXHIBIT A

Fischer sequence

IVALPXGMLK

SEQ ID NO: 2 (single letter sequence)

YFPPPAAKEDFLGCLVKEIPPRLLYAKSSPAYPSVLGQTIRNSRWSSPDNVKPIYIVTPTNASHIQSAVVC GRRHGVRIRVRSGGHDYEGLSYRSLQPEEFAVVDLSKMRAVWVDGKARTAWVDSGAQLGELYYAIHKASTV LAFPAGVCPTIGVGGNFAGGGFGMLLRKYGIAAENVIDVKLVDANGTLHDKKSMGDDHFWAVRGGGGESFG IVVAWKVRLLPVPPTVTVFKIPKKASEGAVDIINRWQVVAPQLPDDLMIRVIAQGPTATFEAMYLGTCQTL TPMMSSKFPELGMNASHCNEMSWIQSIPFVHLGHRDNIEDDLLNRNNTFKPFAEYKSDYVYEPFPKRVWEQ IFSTWLLKPGAGIMIFDPYGATISATPEWATPFPHRKGVLFNIQYVNYWFAPGAGAAPLSWSKEIYNYMEP YVSKNPRQAYANYRDIDLGRNEVVNDVSTFSSGLVWGQKYFKGNFQRLAITKGKVDPTDYFRNEQSIPPLIKKY

SIM+LALNVIEW analysis

37.5% identity in 8 residues overlap; Score: 22.0; Gap frequency: 0.0%

Fischer, 1 IVALPXGM Phlp4(#2), 142 VLAFPAGV * * *

SEQ ID NO: 4 (single letter sequence)

YFPPPAAKEDFLGCLVKEIPPRLLYAKSSPAYPSVLGQTIRNSRWSSPDNVKPIYIVTPTNASHIQSAVVC GRRHGVRIRVRSGGHDYEGLSYRSLQPEEFAVVDLSKMRAVWVDGKARTAWVDSGAQLGELYYAIHKASPV LAFPAGVCPTIGVGGNFAGGGFGMLLRKYGIAAENVIDVKLVDANGTLHDKKSMGDDHFWAVRGGGGESFG IVVAWKVRLLPVPPTVTVFKIPKKASEGAVDIINRWQVVAPQLPDDLMIRVIAQGPTATFEAMYLGTCQTL TPMMSSKFPELGMNASHCNEMSWIQSIPFVHLGHRDNIEDDLLNRNNTFKPFAEYKSDYVYEPFPKEVWEQ IFSTWLLKPGAGIMIFDPYGATISATPEWATPFPHRKGVLFNIQYVNYWFAPGAGAAPLSWSKEIYNYMEP YVSKNPRQAYANYRDIDLGRNEVVNDVSTFSSGLVWGQKYFKGNFQRLAITKGKVDPTDYFRNEQSIPPLIKKY

SIM+LALNVIEW analysis

37.5% identity in 8 residues overlap; Score: 22.0; Gap frequency: 0.0%

Fischer, 1 IVALPXGM Phlp4(#4), 142 VLAFPAGV * * *

SEQ ID NO: 6 (single letter sequence)

YFPPPAAKEDFLGCLVKEIPPRLLYAKSSPAYPSVLGQTIRNSRWSSPDNVKPLYIITPTNVSHIQSAVVC GRRHSVRIRVRSGGHDYEGLSYRSLQPETFAVVDLNKMRAVWVDGKARTAWVDSGAQLGELYYAIYKASPT LAFPAGVCPTIGVGGNFAGGGFGMLLRKYGIAAENVIDVKLVDANGKLHDKKSMGDDHFWAVRGGGGESFG IVVAWQVKLLPVPPTVTIFKISKTVSEGAVDIINKWQVVAPQLPADLMIRIIAQGPKATFEAMYLGTCKTL TPLMSSKFPELGMNPSHCNEMSWIQSIPFVHLGHRDALEDDLLNRNNSFKPFAEYKSDYVYQPFPKTVWEQ ILNTWLVKPGAGIMIFDPYGATISATPESATPFPHRKGVLFNIQYVNYWFAPGAAAAPLSWSKDIYNYMEP YVSKNPRQAYANYRDIDLGRNEVVNDVSTYASGKVWGQKYFKGNFERLAITKGKVDPTDYFRNEQSIPPLI KKY

SIM+LALNVIEW analysis

42.9% identity in 7 residues overlap; Score: 19.0; Gap frequency: 0.0%

Fischer, 2 VALPXGM Phlp4(#6), 143 LAFPAGV * * *